AMENDMENTS TO THE SPECIFICATION

Please delete the paragraph on page 10, line 31, to page 11, line 14, and replace it with the following amended paragraph:

FIGURE 7. Sequence fDEC-205. (A) Schematic representation of DEC-205. (B) The predicted amino acid sequence of DEC-205 (SEQ ID NO:3) is aligned with the sequences of the bovine PLA2 receptor (SEQ ID NO:4) and the human macrophage mannose receptor (SEQ ID NO:5). Amino acid positions where there is identity among all three proteins are shaded. Protein domains are separated, and consensus amino acids that define C-type CRDs (Weis et al., Science 254:1608-15) are indicated blow the relevant sequence as follows: invariant amino acids are shown in single letter code, θ = aliphatic, χ = aliphatic or aromatic, Φ = aromatic, Z= E or Q, B = D or N, O = D, N, E, or Q. The two missing cysteines in CRD 8 are highlighted with a *. **The consensus sequences 'ΦθGθΩΩ', 'EΩCθ ', 'ΦθGθ', 'ZPBB', 'ΦθGθΩ' and 'EΩCθχ' are disclosed as SEQ ID NOS 7-12, respectively.** Peptide sequences determined by automated Edman degradation from purified DEC-205 protein are overlined and numbered (N indicates amino terminal, T ~~indicates~~ **indicated** peptides generated with Trypsin, and L indicates peptides generated with endoproteinase lys-C). (C) Comparison of carboxyl-terminal cytoplasmic domain sequences of human (*top*) (SEQ ID NO:1) and murine (*bottom*) (SEQ ID NO:6) DEC-205. Regions of identity are underlined; regions of similarity are italicized.